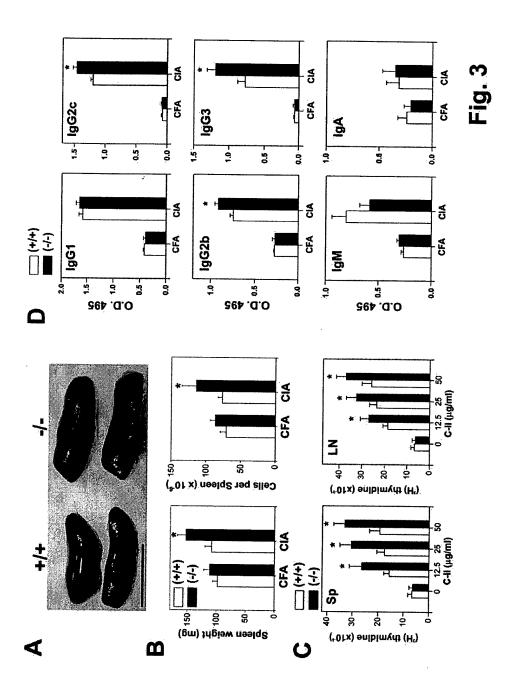
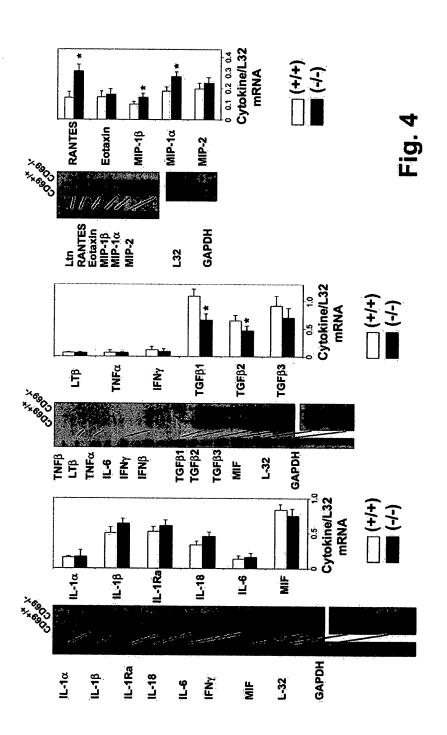
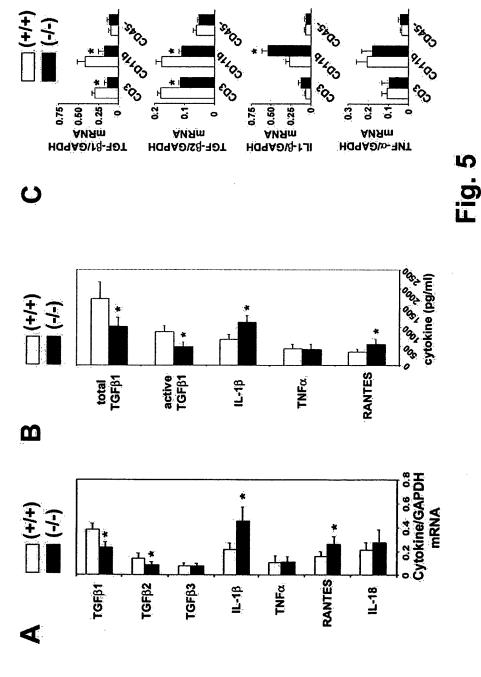
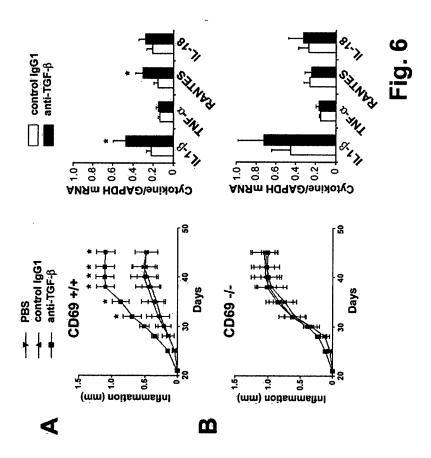


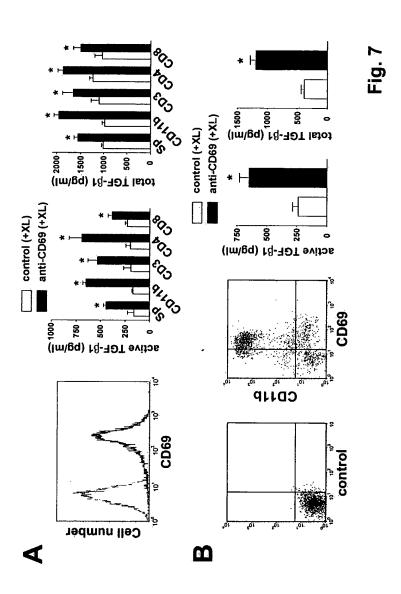
Fig. 2

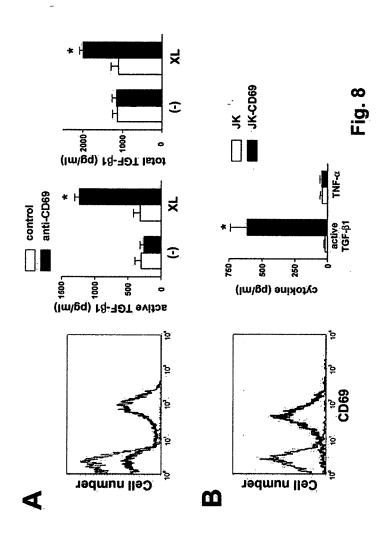


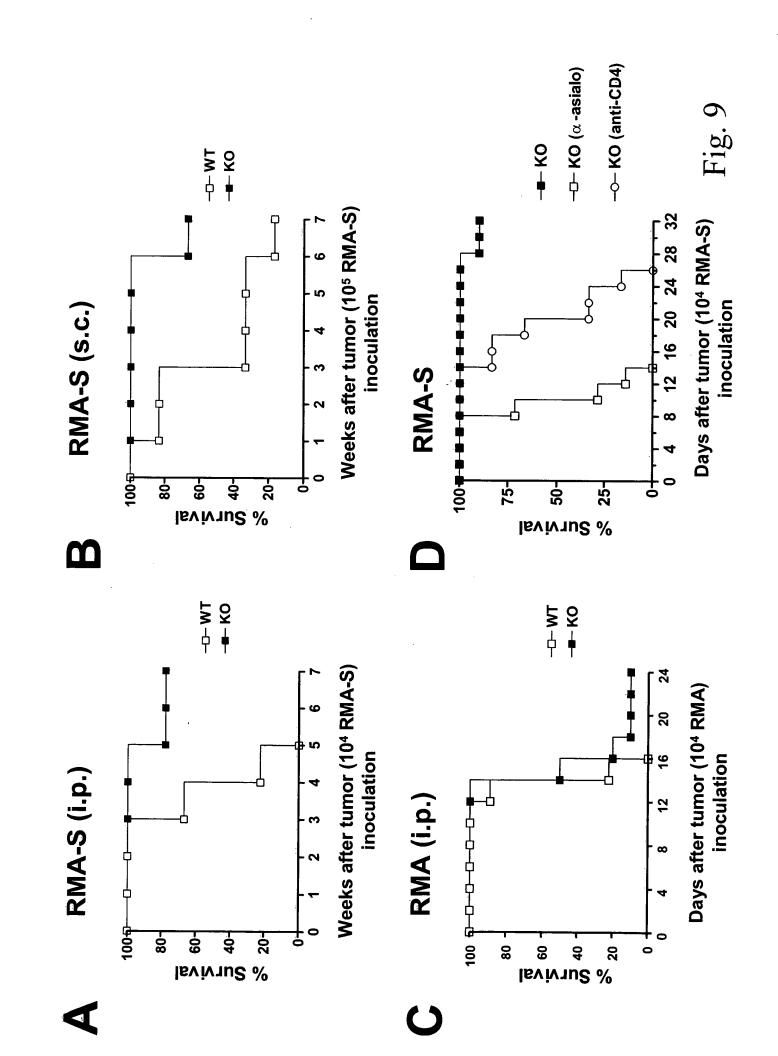








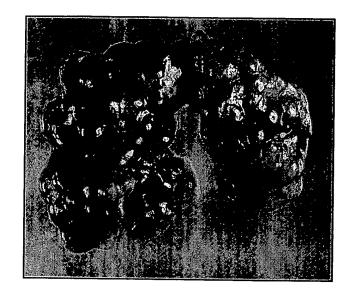




E

CD69 +/+







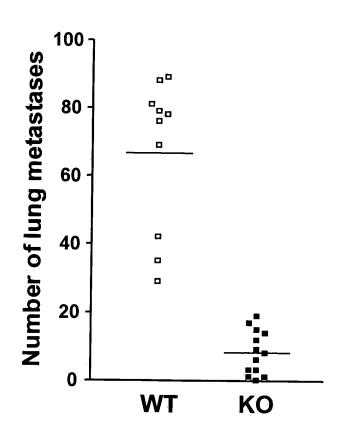
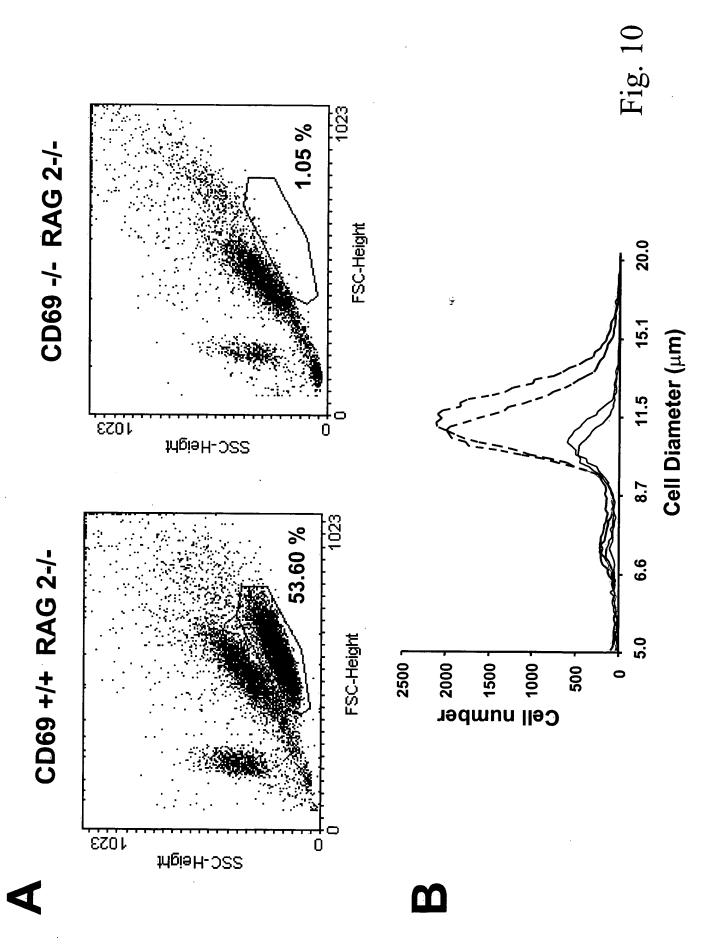


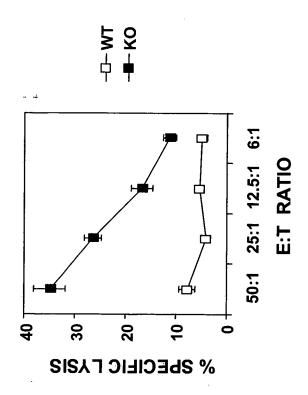
Fig Q

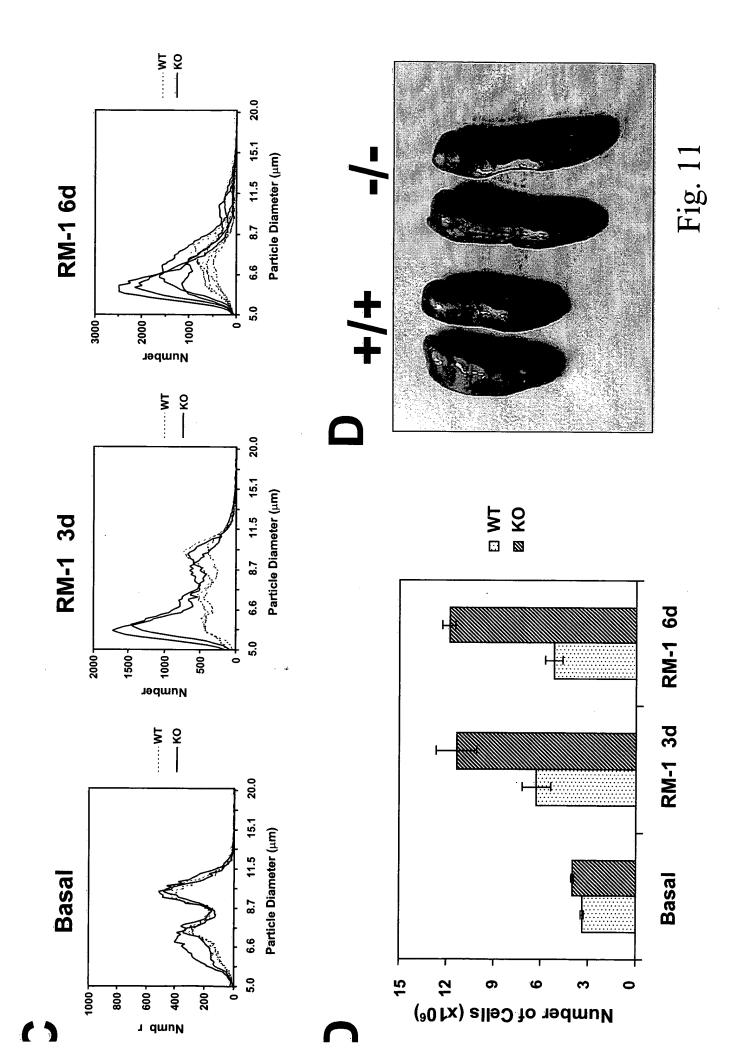


SPECIFIC LYSIS
% PECIFIC LYSIS
% FET RATIO

Fig. 11

മ





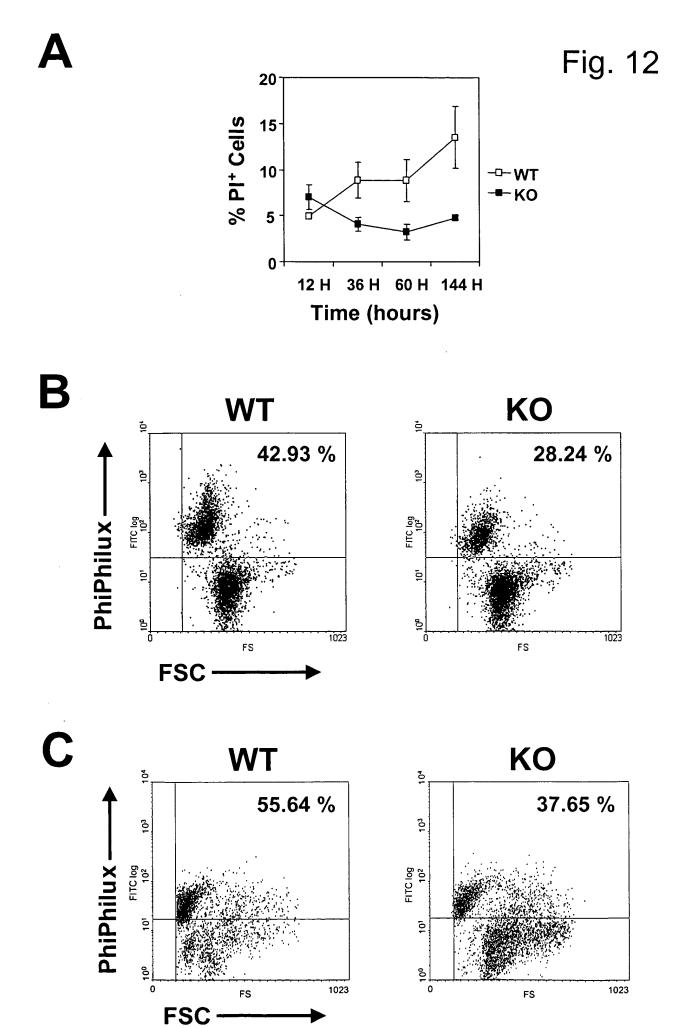
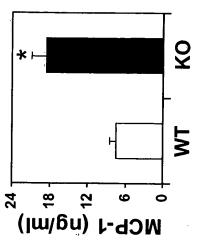
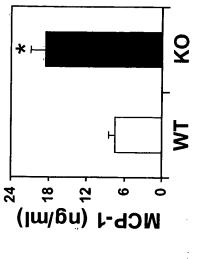


Fig. 13





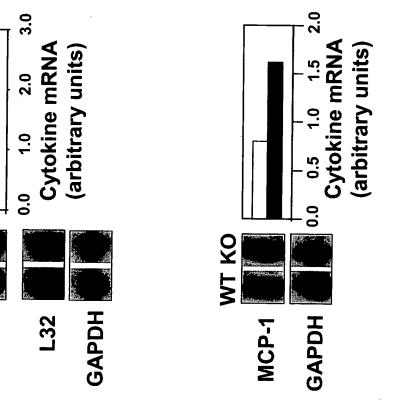


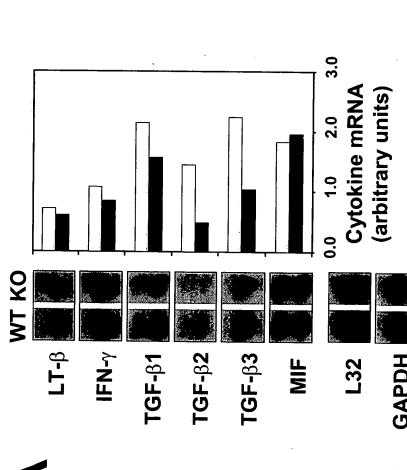
.5

Cytokine mRNA (arbitrary units)

L32

GAPDH





IL 18

IL 6

IL-1 RA

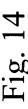
WT KO

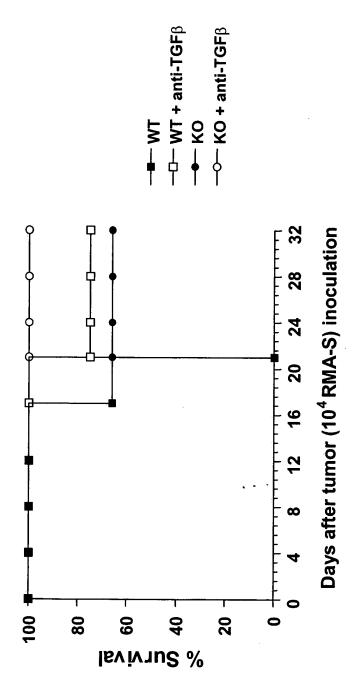
IL 12 p35

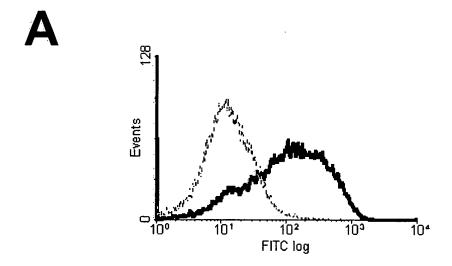
IL-1α

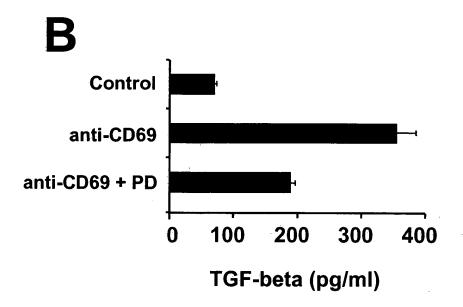
L-1β

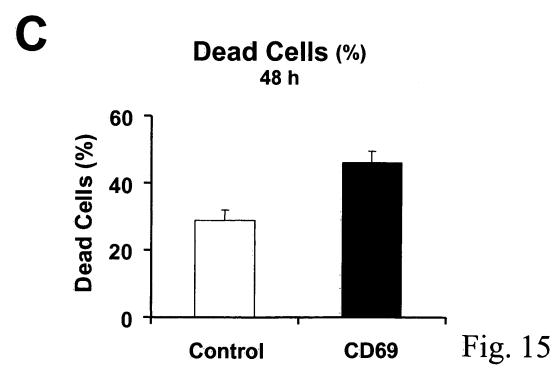
ż

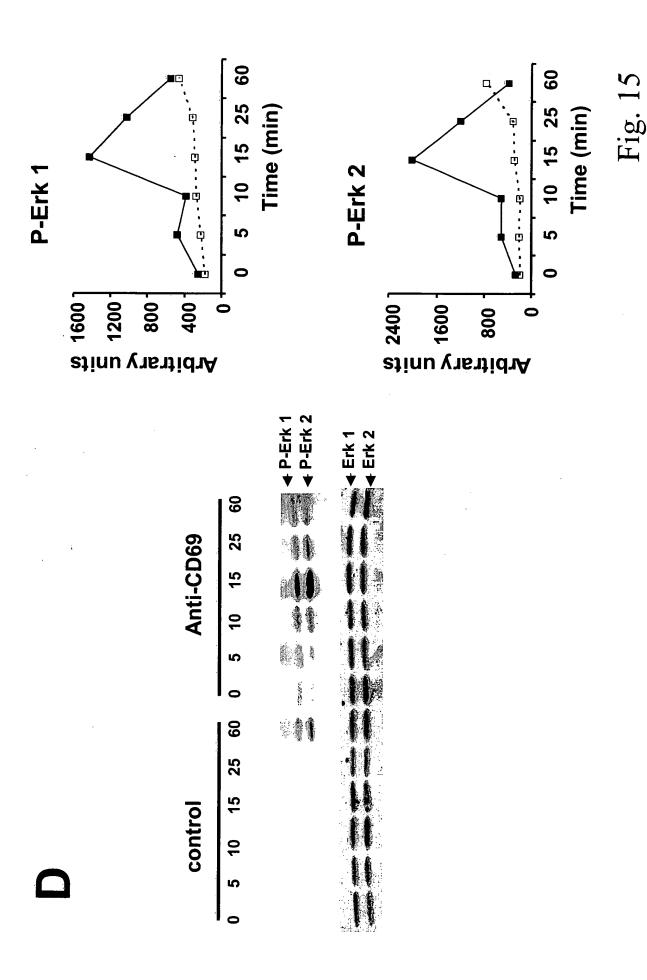






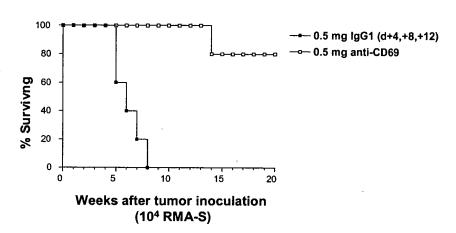






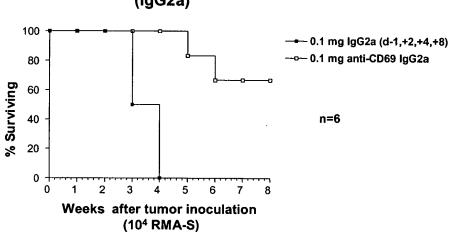
anti-CD69 treatment

(lgG1)



anti-CD69 treatment

(lgG2a)



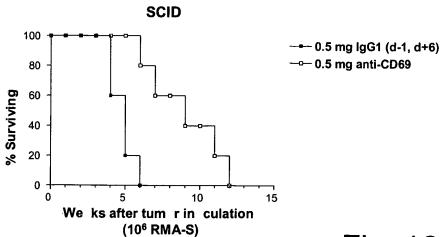
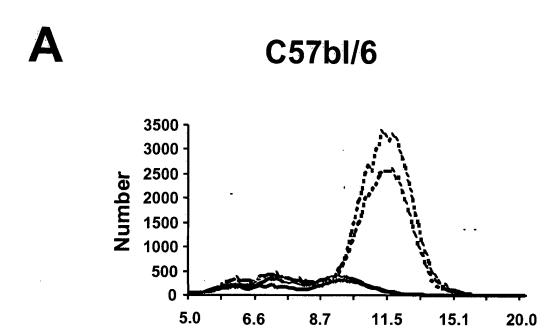


Fig. 16



Particle Diameter (µm)

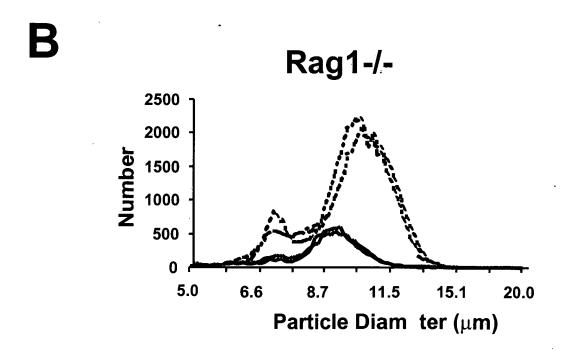
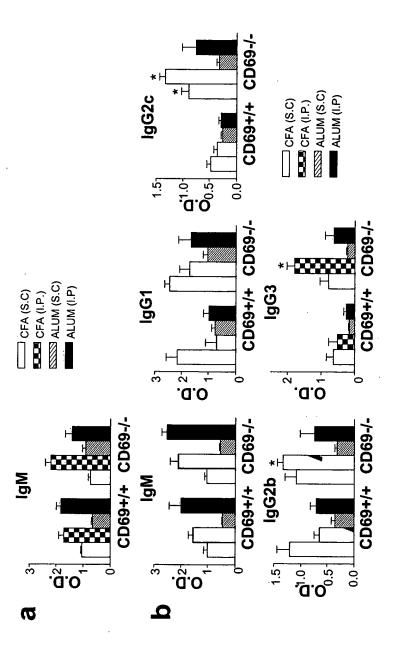
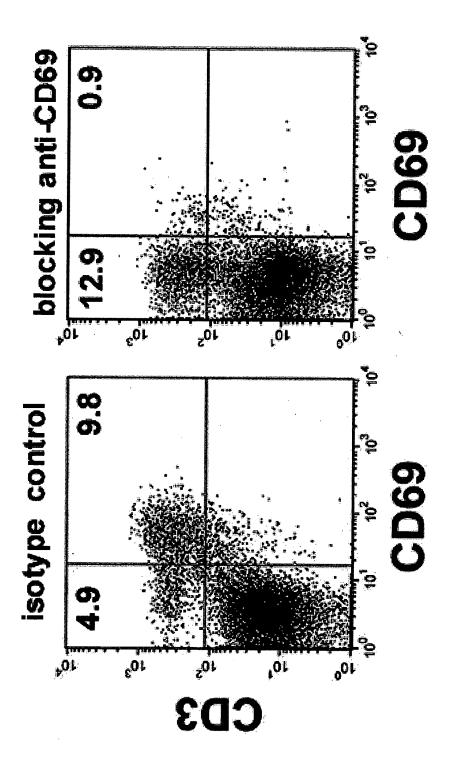
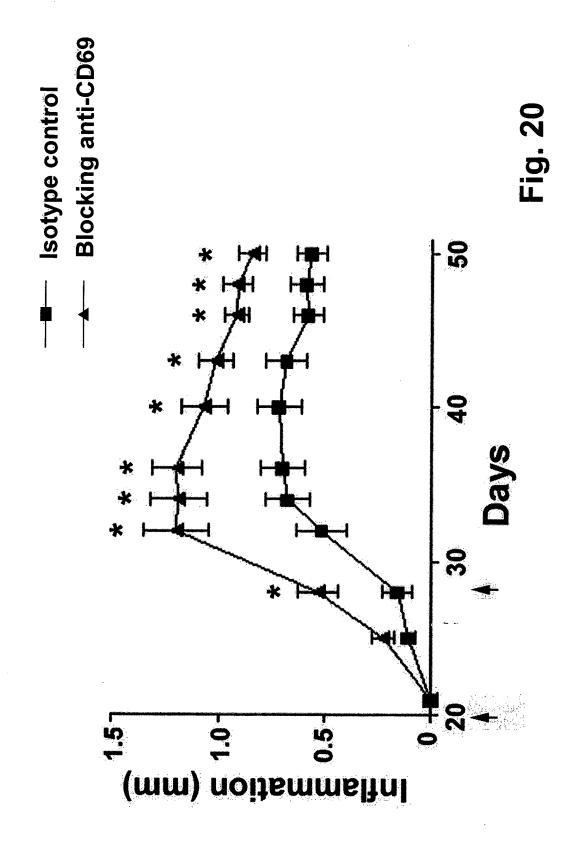


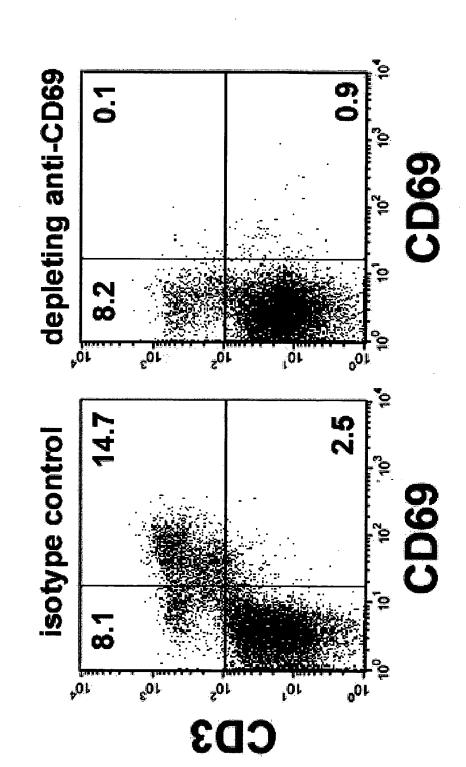
Fig. 17







CIA treatment with depleting anti-CD69



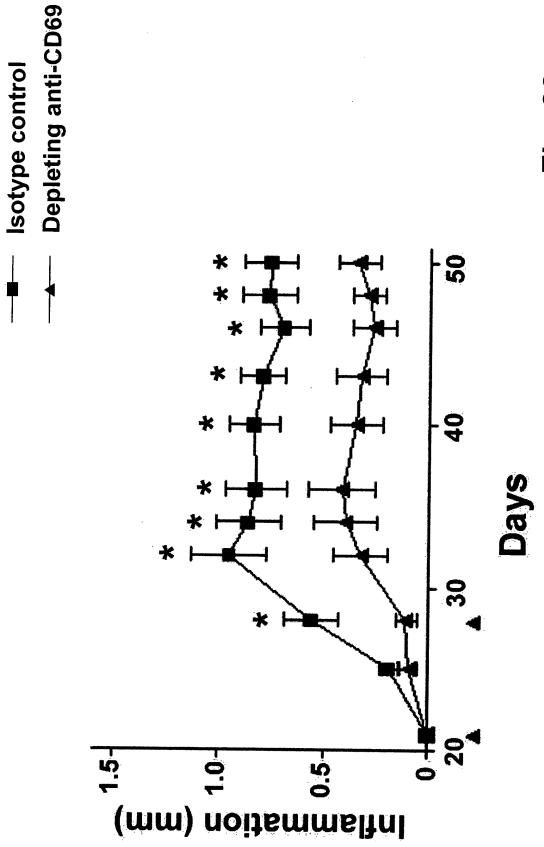


Figure 23.

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(SEQ ID NO:2)
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1 agactcaaca agagctccag caaagacttt cactgtagct tgacttgacc tgagattaac 61 tagggaatct tgagaataaa gatgagctct gaaaattgtt tcgtagcaga gaacagctct 121 ttgcatccgg agagtggaca agaaaatgat gccaccagtc cccatttctc aacacgtcat 181 gaagggtcct tccaagttcc tgtcctgtgt gctgtaatga atgtggtctt catcaccatt 241 ttaatcatag ctctcattgc cttatcagtg ggccaataca attgtccagg ccaatacaca 301 ttctcaatgc catcagacag ccatgtttct tcatgctctg aggactgggt tggctaccag 361 aggaaatgct actttatttc tactgtgaag aggagctgga cttcagccca aaatgcttgt 421 tetgaacatg gtgetaetet tgetgteatt gattetgaaa aggacatgaa etttetaaaa 481 cgatacgcag gtagagaga acactgggtt ggactgaaaa aggaacctgg tcacccatgg 541 aagtggtcaa atggcaaaga atttaacaac tggttcaacg ttacagggtc tgacaagtgt 601 gtttttctga aaaacacaga ggtcagcagc atggaatgtg agaagaattt atactggata 661 tgtaacaaac cttacaaata ataaggaaac atgttcactt attgactatt atagaatgga 721 actcaaggaa atctgtgtca gtggatgctg ctctgtggtc cgaagtcttc catagagact 781 ttgtgaaaaa aaattttata gtgtcttggg aattttcttc caaacagaac tatggaaaaa 841 aaggaagaaa ttccaggaaa atctgcactg tgggctttta ttgccatgag ctagaagcat 901 cacaggttga ccaataacca tgcccaagaa tgagaagaat gactatgcaa cctttggatg 961 cactttatat tattttgaat ccagaaataa tgaaataact aggcgtggac ttactattta 1021 ttgctgaatg actaccaaca gtgagagccc ttcatqcatt tqcactactg gaaggagtta 1081 gatgttggta ctagatactg aatgtaaaca aaggaattat qqctqqtaac ataqqttttt 1141 agtctaattg aatcccttaa actcagggag catttataaa tggacaaatg cttatqaaac 1201 taagatttgt aatatttctc tctttttaga gaaatttgcc aatttacttt gttatttttc 1261 cccaaaaaga atgggatgat cgtgtattta tttttttact tcctcagctg tagacaggtc 1321 cttttcgatg gtacatattt ctttgccttt ataatctttt atacagtgtc ttacagagaa 1381 aagacataag caaagactat gaggaatatt tgcaagacat agaatagtgt tggaaaatgt 1441 gcaatatgtg atgtggcaaa tctctattag gaaatattct gtaatcttca gacctagaat 1501 aatactagtc ttataatagg tttgtgactt tcctaaatca attctattac gtgcaatact 1561 tcaatacttc atttaaaata tttttatgtg caataaaatg tatttgtttg tattttgtgt 1621 tcagtacaat tataagctgt ttttatatat gtgaaataaa agtagaataa acacaaaaaa 1681 aaaaaaaaaa aaaaaaaaa aa

(SEQ ID NO:1)

Figure 24.

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```

Figure 25.

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1 mhdsnnvekd itpselpanp gclhskehsi katliwrlff limfltiivc gmvaalsair
61 anchqepsvc lqaacpeswi gfqrkcfyfs ddtknwtssq rfcdsqdadl aqvesfqeln
121 fllrykgpsd hwiglsreqg qpwkwingte wtrqfpilga gecaylndkg assarhyter
181 kwicsksdih v
(SEQ ID NO:6)
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1 gaattccggc aaaatgcatg acagtaacaa tgtggagaaa gacattacac catctgaatt
 61 gcctgcaaac ccaggttgtc tgcattcaaa agagcattct attaaagcta ccttaatttg
121 gegettattt ttettaatea tgtttetgae aateatagtg tgtggaatgg ttgetgettt
181 aagcgcaata agagctaact gccatcaaga gccatcagta tgtcttcaag ctgcatgccc
241 agaaagctgg attggttttc aaagaaagtg tttctatttt tctgatgaca ccaagaactg
301 gacatcaagt cagaggtttt gtgactcaca agatgctgat cttgctcagg ttgaaagctt
361 ccaggaactg aatttcctgt tgagatataa aggcccatct gatcactgga ttgggctgag
421 cagagaacaa ggccaaccat ggaaatggat aaatggtact gaatggacaa gacagtttcc
481 tatcctggga gcaggagagt gtgcctattt gaatgacaaa ggtgccagta gtgccaggca
541 ctacacagag aggaagtgga tttgttccaa atcagatata catgtctaga tgttacagca
601 aagccccaac taatctttag aagcatattg gaactgataa ctccatttta aaatgagcaa
661 agaatttatt tottatacca acaggtatat gaaaatatgo toaatatcac taataactgg
721 gaaaatacaa atcaaaatca tagtaaaata ttacctgttt tcatggtgct aatattacct
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841 ttgaaaaaa
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TRA 1875815v1